

Title: Inbred Maize Line PH48V

Anticipated Class: 800

Subclass: 200

Art Unit: 1649

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Field of the Invention

This invention is in the field of maize breeding, specifically relating to an inbred maize line designated PH48V.

Background of the Invention

The goal of plant breeding is to combine in a single variety or hybrid various desirable traits. For field crops, these traits may include resistance to diseases and insects, tolerance to heat and drought, reducing the time to crop maturity, greater yield, and better agronomic quality. With mechanical harvesting of many crops, uniformity of plant characteristics such as germination and stand establishment, growth rate, maturity, and plant and ear height, is important.

Field crops are bred through techniques that take advantage of the plant's method of pollination. A plant is self-pollinated if pollen from one flower is transferred to the same or another flower of the same plant. A plant is cross-pollinated if the pollen comes from a flower on a different plant.

Plants that have been self-pollinated and selected for type for many generations become homozygous at almost all gene loci and produce a uniform population of true breeding progeny. A cross between two different homozygous lines produces a uniform population of hybrid plants that may be heterozygous for many gene loci. A cross of two plants each heterozygous at a number of gene loci will produce a population of hybrid plants that differ genetically and will not be uniform.

Maize (*zea mays* L.), often referred to as corn in the United States, can be bred by both self-pollination and cross-pollination techniques. Maize has separate male and female flowers on the same plant, located on the tassel and the ear, respectively. Natural pollination occurs in maize when wind blows pollen from the tassels to the silks that protrude from the tops of the ears.

A reliable method of controlling male fertility in plants offers the opportunity for improved plant breeding. This is especially true for development of maize hybrids,

which relies upon some sort of male sterility system. There are several options for controlling male fertility available to breeders, such as: manual or mechanical emasculation (or detasseling), cytoplasmic male sterility, genetic male sterility, gametocides and the like.

5 Hybrid maize seed is typically produced by a male sterility system incorporating manual or mechanical detasseling. Alternate strips of two maize inbreds are planted in a field, and the pollen-bearing tassels are removed from one of the inbreds (female). Providing that there is sufficient isolation from sources of foreign maize pollen, the ears of the detasseled inbred will be fertilized only from the other inbred (male), and the
10 resulting seed is therefore hybrid and will form hybrid plants.

 The laborious, and occasionally unreliable, detasseling process can be avoided by using cytoplasmic male-sterile (CMS) inbreds. Plants of a CMS inbred are male sterile as a result of factors resulting from the cytoplasmic, as opposed to the nuclear, genome. Thus, this characteristic is inherited exclusively through the female parent in
15 maize plants, since only the female provides cytoplasm to the fertilized seed. CMS plants are fertilized with pollen from another inbred that is not male-sterile. Pollen from the second inbred may or may not contribute genes that make the hybrid plants male-fertile. Seed from detasseled fertile maize and CMS produced seed of the same hybrid can be blended to insure that adequate pollen loads are available for fertilization when
20 the hybrid plants are grown.

 There are several methods of conferring genetic male sterility available, such as multiple mutant genes at separate locations within the genome that confer male sterility, as disclosed in U.S. Patents 4,654,465 and 4,727,219 to Brar et al. and chromosomal translocations as described by Patterson in U.S. Patents No. 3,861,709 and 3,710,511.
25 These and all patents referred to are incorporated by reference. In addition to these methods, Albertsen et al., of Pioneer Hi-Bred, U.S. Patent No. 5,432,068, have developed a system of nuclear male sterility which includes: identifying a gene which is critical to male fertility; silencing this native gene which is critical to male fertility; removing the native promoter from the essential male fertility gene and replacing it with
30 an inducible promoter; inserting this genetically engineered gene back into the plant; and thus creating a plant that is male sterile because the inducible promoter is not "on" resulting in the male fertility gene not being transcribed. Fertility is restored by inducing, or turning "on", the promoter, which in turn allows the gene that confers male fertility to be transcribed.

There are many other methods of conferring genetic male sterility in the art, each with its own benefits and drawbacks. These methods use a variety of approaches such as delivering into the plant a gene encoding a cytotoxic substance associated with a male tissue specific promoter or an antisense system in which a gene critical to fertility is identified and an antisense to that gene is inserted in the plant (see: Fabinjanski, et al. EPO 89/3010153.8 publication no. 329,308 and PCT application PCT/CA90/00037 published as WO 90/08828).

Another system useful in controlling male sterility makes use of gametocides. Gametocides are not a genetic system, but rather a topical application of chemicals. These chemicals affect cells that are critical to male fertility. The application of these chemicals affects fertility in the plants only for the growing season in which the gametocide is applied (see Carlson, Glenn R., U.S. Patent Number: 4,936,904). Application of the gametocide, timing of the application and genotype specificity often limit the usefulness of the approach.

Development of Maize Inbred Lines

The use of male sterile inbreds is but one factor in the production of maize hybrids. Plant breeding techniques known in the art and used in a maize plant breeding program include, but are not limited to, recurrent selection, backcrossing, pedigree breeding, restriction length polymorphism enhanced selection, genetic marker enhanced selection and transformation. The development of maize hybrids in a maize plant breeding program requires, in general, the development of homozygous inbred lines, the crossing of these lines, and the evaluation of the crosses. Pedigree breeding and recurrent selection breeding methods are used to develop inbred lines from breeding populations. Maize plant breeding programs combine the genetic backgrounds from two or more inbred lines or various other germplasm sources into breeding pools from which new inbred lines are developed by selfing and selection of desired phenotypes. The new inbreds are crossed with other inbred lines and the hybrids from these crosses are evaluated to determine which of those have commercial potential. Plant breeding and hybrid development, as practiced in a maize plant breeding program, are expensive and time consuming processes.

Pedigree breeding starts with the crossing of two genotypes, each of which may have one or more desirable characteristics that is lacking in the other or which complements the other. If the two original parents do not provide all the desired characteristics, other sources can be included in the breeding population. In the

pedigree method, superior plants are selfed and selected in successive generations. In the succeeding generations the heterozygous condition gives way to homogeneous lines as a result of self-pollination and selection. Typically in the pedigree method of breeding five or more generations of selfing and selection is practiced: $F_1 \rightarrow F_2$; $F_2 \rightarrow F_3$;

5 $F_3 \rightarrow F_4$; $F_4 \rightarrow F_5$, etc.

Recurrent selection breeding, backcrossing for example, can be used to improve an inbred line and a hybrid which is made using those inbreds. Backcrossing can be used to transfer a specific desirable trait from one inbred or source to an inbred that lacks that trait. This can be accomplished, for example, by first crossing a superior
10 inbred (recurrent parent) to a donor inbred (non-recurrent parent), that carries the appropriate gene(s) for the trait in question. The progeny of this cross is then mated back to the superior recurrent parent followed by selection in the resultant progeny for the desired trait to be transferred from the non-recurrent parent. After five or more
15 backcross generations with selection for the desired trait, the progeny will be homozygous for loci controlling the characteristic being transferred, but will be like the superior parent for essentially all other genes. The last backcross generation is then selfed to give pure breeding progeny for the gene(s) being transferred. A hybrid developed from inbreds containing the transferred gene(s) is essentially the same as a hybrid developed from the same inbreds without the transferred gene(s).

20 Elite inbred lines, that is, pure breeding, homozygous inbred lines, can also be used as starting materials for breeding or source populations from which to develop other inbred lines. These inbred lines derived from elite inbred lines can be developed using the pedigree breeding and recurrent selection breeding methods described earlier. As an example, when backcross breeding is used to create these derived lines in a
25 maize plant breeding program, elite inbreds can be used as a parental line or starting material or source population and can serve as either the donor or recurrent parent.

Development of Maize Hybrids

A single cross maize hybrid results from the cross of two inbred lines, each of which has a genotype that complements the genotype of the other. The hybrid progeny
30 of the first generation is designated F_1 . In the development of commercial hybrids in a maize plant breeding program, only the F_1 hybrid plants are sought. Preferred F_1 hybrids are more vigorous than their inbred parents. This hybrid vigor, or heterosis, can be manifested in many polygenic traits, including increased vegetative growth and increased yield.

The development of a maize hybrid in a maize plant breeding program involves three steps: (1) the selection of plants from various germplasm pools for initial breeding crosses; (2) the selfing of the selected plants from the breeding crosses for several generations to produce a series of inbred lines, which, although different from each other, breed true and are highly uniform; and (3) crossing the selected inbred lines with different inbred lines to produce the hybrid progeny (F_1). During the inbreeding process in maize, the vigor of the lines decreases. Vigor is restored when two different inbred lines are crossed to produce the hybrid progeny (F_1). An important consequence of the homozygosity and homogeneity of the inbred lines is that the hybrid between a defined pair of inbreds will always be the same. Once the inbreds that give a superior hybrid have been identified, the hybrid seed can be reproduced indefinitely as long as the homogeneity of the inbred parents is maintained.

A single cross hybrid is produced when two inbred lines are crossed to produce the F_1 progeny. A double cross hybrid is produced from four inbred lines (or synthetics) crossed in pairs ($A \times B$ and $C \times D$) and then the two F_1 hybrids are crossed again ($A \times B$) \times ($C \times D$). A three-way cross hybrid is produced from three inbred lines (or synthetics) where two of the inbred lines (or synthetics) are crossed ($A \times B$) and then the resulting F_1 hybrid is crossed with the third inbred (or synthetics) ($A \times B$) \times C . Much of the hybrid vigor exhibited by F_1 hybrids is lost in the next generation (F_2). Consequently, seed from hybrids is not used for planting stock.

Hybrid seed production requires elimination or inactivation of pollen produced by the female parent. Incomplete removal or inactivation of the pollen provides the potential for self pollination. This inadvertently self pollinated seed may be unintentionally harvested and packaged with hybrid seed.

Once the seed is planted, it is possible to identify and select these self pollinated plants. These self pollinated plants will be genetically equivalent to the female inbred line used to produce the hybrid.

Typically these self pollinated plants can be identified and selected due to their decreased vigor. Female selfs are identified by their less vigorous appearance for vegetative and/or reproductive characteristics, including shorter plant height, small ear size, ear and kernel shape, cob color, or other characteristics.

Identification of these self-pollinated lines can also be accomplished through molecular marker analyses. See, "The Identification of Female Selfs in Hybrid Maize: A Comparison Using Electrophoresis and Morphology", Smith, J.S.C. and Wych, R.D.,

breeder of ordinary skill in the art cannot predict the genotype, how that genotype will interact with various climatic conditions or the resulting phenotypes of the developing lines, except perhaps in a very broad and general fashion. A breeder of ordinary skill in the art would also be unable to recreate the same line twice from the very same original
5 parents as the breeder is unable to direct how the genomes combine or how they will interact with the environmental conditions. This unpredictability results in the expenditure of large amounts of research resources in the development of a superior new maize inbred line.

10 **Summary of the Invention**

According to the invention, there is provided a novel inbred maize line, designated PH48V. This invention thus relates to the seeds of inbred maize line PH48V, to the plants of inbred maize line PH48V, to methods for producing a maize
15 plant produced by crossing the inbred maize line PH48V with itself or another maize line, and to methods for producing a maize plant containing in its genetic material one or more transgenes and to the transgenic maize plants produced by that method. This invention also relates to inbred maize lines derived from inbred maize line PH48V, to methods for producing other inbred maize lines derived from inbred maize line PH48V and to the inbred maize lines derived by the use of those methods. This invention
20 further relates to hybrid maize seeds and plants produced by crossing the inbred line PH48V with another maize line.

Definitions

In the description and examples that follow, a number of terms are used herein.
25 In order to provide a clear and consistent understanding of the specification and claims, including the scope to be given such terms, the following definitions are provided. NOTE: ABS is in absolute terms and %MN is percent of the mean for the experiments in which the inbred or hybrid was grown. These designators will follow the descriptors to denote how the values are to be interpreted. Below are the descriptors used in the data
30 tables included herein.

ANT ROT = ANTHRACNOSE STALK ROT (*Colletotrichum graminicola*). A 1 to 9 visual rating indicating the resistance to Anthracnose Stalk Rot. A higher score indicates a higher resistance.

5 GDU SHD = GDU TO SHED. The number of growing degree units (GDUs) or heat units required for an inbred line or hybrid to have approximately 50 percent of the plants shedding pollen and is measured from the time of planting. Growing degree units are calculated by the Barger Method, where the heat units for a 24-hour period are:

$$\text{GDU} = \frac{(\text{Max. temp.} + \text{Min. temp.})}{2} - 50$$

10 The highest maximum temperature used is 86° F. and the lowest minimum temperature used is 50° F. For each inbred or hybrid it takes a certain number of GDUs to reach various stages of plant development.

GDU SLK = GDU TO SILK. The number of growing degree units required for an inbred line or hybrid to have approximately 50 percent of the plants with silk emergence from time of planting. Growing degree units are calculated by the Barger Method as given in GDU SHD definition.

15 GIB ERS = GIBBERELLA EAR ROT (PINK MOLD) (*Gibberella zeae*). A 1 to 9 visual rating indicating the resistance to Gibberella Ear Rot. A higher score indicates a higher resistance.

20 GLF SPT = Gray Leaf Spot (*Cercospora zeae-maydis*). A 1 to 9 visual rating indicating the resistance to Gray Leaf Spot. A higher score indicates a higher resistance.

GOS WLT = Goss' Wilt (*Corynebacterium nebraskense*). A 1 to 9 visual rating indicating the resistance to Goss' Wilt. A higher score indicates a higher resistance.

25 GRN APP = GRAIN APPEARANCE. This is a 1 to 9 rating for the general appearance of the shelled grain as it is harvested based on such factors as the color of harvested grain, any mold on the grain, and any cracked grain. High scores indicate good grain quality.

30 H/POP = YIELD AT HIGH DENSITY. Yield ability at relatively high plant densities on 1-9 relative rating system with a higher number indicating the hybrid responds well to high plant densities for yield relative to other hybrids. A 1, 5, and 9 would represent very poor, average, and very good yield response, respectively, to increased plant density.

HC BLT = HELMINTHOSPORIUM CARBONUM LEAF BLIGHT (*Helminthosporium carbonum*). A 1 to 9 visual rating indicating the resistance to Helminthosporium infection. A higher score indicates a higher resistance.

PLT HT = PLANT HEIGHT. This is a measure of the height of the plant from the ground to the tip of the tassel in inches.

POL SC = POLLEN SCORE. A 1 to 9 visual rating indicating the amount of pollen shed. The higher the score the more pollen shed.

5 POL WT = POLLEN WEIGHT. This is calculated by dry weight of tassels collected as shedding commences minus dry weight from similar tassels harvested after shedding is complete.

POP K/A = PLANT POPULATIONS. Measured as 1000s per acre.

10 POP ADV = PLANT POPULATION ADVANTAGE. The plant population advantage of variety #1 over variety #2 as calculated by PLANT POPULATION of variety #2 - PLANT POPULATION of variety #1 = PLANT POPULATION ADVANTAGE of variety #1.

15 PRM = PREDICTED RELATIVE MATURITY. This trait, predicted relative maturity, is based on the harvest moisture of the grain. The relative maturity rating is based on a known set of checks and utilizes standard linear regression analyses and is also referred to as the Comparative Relative Maturity Rating System that is similar to the Minnesota Relative Maturity Rating System.

20 PRM SHD = A relative measure of the growing degree units (GDU) required to reach 50% pollen shed. Relative values are predicted values from the linear regression of observed GDU's on relative maturity of commercial checks.

PROT = GRAIN PROTEIN. Absolute value of protein content of the kernel as predicted by Near-Infrared Transmittance and expressed as a percent of dry matter.

25 RT LDG = ROOT LODGING. Root lodging is the percentage of plants that do not root lodge; plants that lean from the vertical axis at an approximately 30° angle or greater would be counted as root lodged.

RTL ADV = ROOT LODGING ADVANTAGE. The root lodging advantage of variety #1 over variety #2.

30 SCT GRN = SCATTER GRAIN. A 1 to 9 visual rating indicating the amount of scatter grain (lack of pollination or kernel abortion) on the ear. The higher the score the less scatter grain.

SDG VGR = SEEDLING VIGOR. This is the visual rating (1 to 9) of the amount of vegetative growth after emergence at the seedling stage (approximately five leaves). A higher score indicates better vigor.

TEX EAR = EAR TEXTURE. A 1 to 9 visual rating was used to indicate the relative hardness (smoothness of crown) of mature grain. A 1 would be very soft (extreme dent) while a 9 would be very hard (flinty or very smooth crown).

TILLER = TILLERS. A count of the number of tillers per plot that could possibly shed pollen was taken. Data are given as a percentage of tillers: number of tillers per plot divided by number of plants per plot.

TST WT = TEST WEIGHT (UNADJUSTED). The measure of the weight of the grain in pounds for a given volume (bushel).

TST WTA = TEST WEIGHT ADJUSTED. The measure of the weight of the grain in pounds for a given volume (bushel) adjusted for 15.5 percent moisture.

TSW ADV = TEST WEIGHT ADVANTAGE. The test weight advantage of variety #1 over variety #2.

WIN M% = PERCENT MOISTURE WINS.

WIN Y% = PERCENT YIELD WINS.

YLD = YIELD. It is the same as BU ACR ABS.

YLD ADV = YIELD ADVANTAGE. The yield advantage of variety #1 over variety #2 as calculated by: YIELD of variety #1 - YIELD variety #2 = yield advantage of variety #1.

YLD SC = YIELD SCORE. A 1 to 9 visual rating was used to give a relative rating for yield based on plot ear piles. The higher the rating the greater visual yield appearance.

Detailed Description of the Invention

Inbred maize lines are typically developed for use in the production of hybrid maize lines. Inbred maize lines need to be highly homogeneous, homozygous and reproducible to be useful as parents of commercial hybrids. There are many analytical methods available to determine the homozygotic and phenotypic stability of these inbred lines.

The oldest and most traditional method of analysis is the observation of phenotypic traits. The data is usually collected in field experiments over the life of the maize plants to be examined. Phenotypic characteristics most often observed are for traits associated with plant morphology, ear and kernel morphology, insect and disease resistance, maturity, and yield.

In addition to phenotypic observations, the genotype of a plant can also be examined. There are many laboratory-based techniques available for the analysis, comparison and characterization of plant genotype; among these are Isozyme Electrophoresis, Restriction Fragment Length Polymorphisms (RFLPs), Randomly Amplified Polymorphic DNAs (RAPDs), Arbitrarily Primed Polymerase Chain Reaction (AP-PCR), DNA Amplification Fingerprinting (DAF), Sequence Characterized Amplified Regions (SCARs), Amplified Fragment Length Polymorphisms (AFLPs), and Simple Sequence Repeats (SSRs) which are also referred to as Microsatellites.

The most widely used of these laboratory techniques are Isozyme Electrophoresis and RFLPs as discussed in Lee, M., "Inbred Lines of Maize and Their Molecular Markers," *The Maize Handbook*, (Springer-Verlag, New York, Inc. 1994, at 423- 432) incorporated herein by reference. Isozyme Electrophoresis is a useful tool in determining genetic composition, although it has relatively low number of available markers and the low number of allelic variants among maize inbreds. RFLPs have the advantage of revealing an exceptionally high degree of allelic variation in maize and the number of available markers is almost limitless.

Maize RFLP linkage maps have been rapidly constructed and widely implemented in genetic studies. One such study is described in Boppenmaier, et al., "Comparisons among strains of inbreds for RFLPs", *Maize Genetics Cooperative Newsletter*, 65:1991, pg. 90, is incorporated herein by reference. This study used 101 RFLP markers to analyze the patterns of 2 to 3 different deposits each of five different inbred lines. The inbred lines had been selfed from 9 to 12 times before being adopted into 2 to 3 different breeding programs. It was results from these 2 to 3 different breeding programs that supplied the different deposits for analysis. These five lines were maintained in the separate breeding programs by selfing or sibbing and rogueing off-type plants for an additional one to eight generations. After the RFLP analysis was completed, it was determined the five lines showed 0-2% residual heterozygosity. Although this was a relatively small study, it can be seen using RFLPs that the lines had been highly homozygous prior to the separate strain maintenance.

Inbred maize line PH48V is a yellow, dent maize inbred that is suited as a female for producing first generation F1 maize hybrids. Inbred maize line PH48V is best adapted to the Southeast region of the United States and can be used to produce hybrids from approximately 121 relative maturity based on the Comparative Relative Maturity Rating System for harvest moisture of grain. Inbred maize line PH48V

demonstrates high resistance to Southern Leaf Blight, high resistance to Northern Leaf Blight and Gray Leaf Spot as an inbred per se. In hybrid combination, including for its area of adaptation, inbred PH48V demonstrates high yield, above average resistance to stalk lodging, above average resistance to root lodging, strong staygreen, tall plant, high ear placement and strong disease resistance.

The inbred has shown uniformity and stability within the limits of environmental influence for all the traits as described in the Variety Description Information (Table 1) that follows. The inbred has been self-pollinated and ear-rowed a sufficient number of generations with careful attention paid to uniformity of plant type to ensure the homozygosity and phenotypic stability necessary to use in commercial production. The line has been increased both by hand and in isolated fields with continued observation for uniformity. No variant traits have been observed or are expected in PH48V.

Inbred maize line PH48V, being substantially homozygous, can be reproduced by planting seeds of the line, growing the resulting maize plants under self-pollinating or sib-pollinating conditions with adequate isolation, and harvesting the resulting seed, using techniques familiar to the agricultural arts.

TABLE 1
VARIETY DESCRIPTION INFORMATION
VARIETY = PH48V

1. TYPE: (describe intermediate types in Comments section):						
5	2	1=Sweet 2=Dent 3=Flint 4=Flour 5=Pop 6=Ornamental				
2. MATURITY:						
	DAYS	HEAT UNITS				
	078	1,534.7	From emergence to 50% of plants in silk			
	076	1,486.7	From emergence to 50% of plants in pollen			
10	002	0,058.3	From 10% to 90% pollen shed			
			From 50% silk to harvest at 25% moisture			
3. PLANT:						
				Standard	Sample	
				Deviation	Size	
15	0,226.7	cm	Plant Height (to tassel tip)	8.14	15	
	0,086.0	cm	Ear Height (to base of top ear node)	7.55	15	
	0,014.3	cm	Length of Top Ear Internode	1.81	15	
	0.0		Average Number of Tillers	0.01	3	
	1.1		Average Number of Ears per Stalk	0.18	3	
	1.0		Anthocyanin of Brace Roots: 1=Absent 2=Faint 3=Moderate 4=Dark			
4. LEAF:						
				Standard	Sample	
				Deviation	Size	
20	010.1	cm	Width of Ear Node Leaf	0.42	15	
	071.6	cm	Length of Ear Node Leaf	1.93	15	
	06.1		Number of leaves above top ear	0.12	15	
25	016.6		Degrees Leaf Angle (measure from 2nd leaf above ear at anthesis to stalk above leaf)	3.02	15	
	03		Leaf Color	Dark Green	(Munsell code)	5GY34
	1.0		Leaf Sheath Pubescence (Rate on scale from 1=none to 9=like peach fuzz)			
			Marginal Waves (Rate on scale from 1=none to 9=many)			
30			Longitudinal Creases (Rate on scale from 1=none to 9=many)			
5. TASSEL:						
				Standard	Sample	
				Deviation	Size	
35	09.9		Number of Primary Lateral Branches	0.31	15	
	034.7		Branch Angle from Central Spike	3.25	15	
	51.5	cm	Tassel Length (from top leaf collar to tassel tip)	1.70	15	
	5.0		Pollen Shed (rate on scale from 0=male sterile to 9=heavy shed)			
	01		Anther Color	Light Green	(Munsell code)	2.5GY88
	01		Glume Color	Light Green	(Munsell code)	7.5GY56
	1.0		Bar Glumes (Glume Bands): 1=Absent 2=Present			
40	22	cm	Peduncle Length (cm. from top leaf to basal branches)			
6a. EAR (Unhusked Data):						
	1	Silk Color (3 days after emergence)	Light Green	(Munsell code)		2.5GY96
	1	Fresh Husk Color (25 days after 50% silking)	Light Green	(Munsell code)		5GY78
	21	Dry Husk Color (65 days after 50% silking)	Buff	(Munsell code)		5Y92
45	1	Position of Ear at Dry Husk Stage: 1= Upright 2= Horizontal 3= Pendant				Upright
	8	Husk Tightness (Rate of Scale from 1=very loose to 9=very tight)				
	3	Husk Extension (at harvest): 1=Short (ears exposed) 2=Medium (<8 cm) 3=Long (8-10 cm beyond ear tip) 4=Very Long (>10 cm)				Long

6b. EAR (Husked Ear Data):		Standard Deviation	Sample Size	
15	cm Ear Length	1.00	15	
33	mm Ear Diameter at mid-point	2.65	15	
47	gm Ear Weight	27.43	15	
17	Number of Kernel Rows	1.15	15	
2	Kernel Rows: 1=Indistinct 2=Distinct			Distinct
2	Row Alignment: 1=Straight 2=Slightly Curved 3=Spiral			Slightly Curved
9	cm Shank Length	1.15	15	
2	Ear Taper: 1=Slight 2= Average 3=Extreme			Average
7. KERNEL (Dried):		Standard Deviation	Sample Size	
9	mm Kernel Length	1.15	15	
8	mm Kernel Width	0.00	15	
7	mmn Kernel Thickness	1.00	15	
88	% Round Kernels (Shape Grade)	5.51	3	
1	Aleurone Color Pattern: 1=Homozygous 2=Segregating			Homozygous
7	Aluerone Color Yellow	(Munsell code)		10YR714
7	Hard Endosperm Color Yellow	(Munsell code)		10YR712
3	Endosperm Type: Normal Starch			
	1=Sweet (Su1) 2=Extra Sweet (sh2) 3=Normal Starch			
	4=High Amylose Starch 5=Waxy Starch 6=High Protein			
	7=High Lysine 8=Super Sweet (se) 9=High Oil			
	10=Other _____			
27	gm Weight per 100 Kernels (unsized sample)	1.53	3	
8. COB:		Standard Deviation	Sample Size	
21	mm Cob Diameter at mid-point	1.15	15	
14	Cob Color Red	(Munsell code)		10R38
9. DISEASE RESISTANCE (Rate from 1 (most susceptible) to 9 (most resistant); leave blank if not tested; leave Race or Strain Options blank if polygenic):				
A. Leaf Blights, Wilts, and Local Infection Diseases				
	Anthracnose Leaf Blight (<i>Colletotrichum graminicola</i>)			
6	Common Rust (<i>Puccinia sorghi</i>)			
	Common Smut (<i>Ustilago maydis</i>)			
	Eyespot (<i>Kabatiella zeae</i>)			
	Goss's Wilt (<i>Clavibacter michiganense</i> spp. <i>nebraskense</i>)			
7	Gray Leaf Spot (<i>Cercospora zeae-maydis</i>)			
	Helminthosporium Leaf Spot (<i>Bipolaris zeicola</i>)	Race _____		
7	Northern Leaf Blight (<i>Exserohilum turcicum</i>)	Race _____		
7	Southern Leaf Blight (<i>Bipolaris maydis</i>)	Race _____		
3	Southern Rust (<i>Puccinia polysora</i>)			
8	Stewart's Wilt (<i>Erwinia stewartii</i>)			
	Other (Specify) _____			
B. Systemic Diseases				
	Corn Lethal Necrosis (MCMV and MDMV)			
	Head Smut (<i>Sphacelotheca reiliana</i>)			
	Maize Chlorotic Dwarf Virus (MDV)			
	Maize Chlorotic Mottle Virus (MCMV)			
4	Maize Dwarf Mosaic Virus (MDMV)			
	Sorghum Downy Mildew of Corn (<i>Peronosclerospora sorghi</i>)			
	Other (Specify) _____			

[illegible]

- #### D. Ear and Kernel Rots

10. INSECT RESISTANCE (Rate from 1 (most susceptible) to 9 (most resistant);
(leave blank if not tested) :

- #### 11. AGRONOMIC TRAITS:

- *In interpreting the foregoing color designations, reference may be made to the Munsell Glossy Book of Color, a standard color reference.

Further Embodiments of the Invention

This invention also is directed to methods for producing a maize plant by crossing a first parent maize plant with a second parent maize plant wherein either the first or second parent maize plant is an inbred maize plant of the line PH48V. Further, both first and second parent maize plants can come from the inbred maize line PH48V. Still further, this invention also is directed to methods for producing an inbred maize line PH48V-derived maize plant by crossing inbred maize line PH48V with a second maize plant and growing the progeny seed, and repeating the crossing and growing steps with the inbred maize line PH48V-derived plant from 0 to 5 times. Thus, any such methods using the inbred maize line PH48V are part of this invention: selfing, backcrosses, hybrid production, crosses to populations, and the like. All plants produced using inbred maize line PH48V as a parent are within the scope of this invention, including plants derived from inbred maize line PH48V. Advantageously, the inbred maize line is used in crosses with other, different, maize inbreds to produce first generation (F₁) maize hybrid seeds and plants with superior characteristics.

A further embodiment of the invention is a single gene conversion or introgression of the maize plant disclosed herein in which the gene or genes of interest (encoding the desired trait) are introduced through traditional (non-transformation) breeding techniques, such as backcrossing (Hallauer *et al*, 1988). One or more genes may be introduced using these techniques. Desired traits transferred through this process include, but are not limited to, waxy starch, nutritional enhancements, industrial enhancements, disease resistance, insect resistance, herbicide resistance and yield enhancements. The gene of interest is transferred from the donor parent to the recurrent parent, in this case, the maize plant disclosed herein. These single gene traits may result from either the transfer of a dominant allele or a recessive allele. Selection of progeny containing the trait of interest is done by direct selection for a trait associated with a dominant allele. Selection of progeny for a trait that is transferred via a recessive allele, such as the waxy starch characteristic, requires growing and selfing the first backcross to determine which plants carry the recessive alleles. Recessive traits may require additional progeny testing in successive backcross generations to determine the presence of the gene of interest.


It should be understood that the inbred can, through routine manipulation of cytoplasmic or other factors, be produced in a male-sterile form. Such embodiments are also contemplated within the scope of the present claims.

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As used herein, the term plant includes plant cells, plant protoplasts, plant cell tissue cultures from which maize plants can be regenerated, plant calli, plant clumps, and plant cells that are intact in plants or parts of plants, such as embryos, pollen, ovules, seeds, flowers, kernels, ears, cobs, leaves, husks, stalks, roots, root tips, 5 anthers, silk and the like.

Duncan, Williams, Zehr, and Widholm, *Planta* (1985) 165:322-332 reflects that 97% of the plants cultured that produced callus were capable of plant regeneration. Subsequent experiments with both inbreds and hybrids produced 91% regenerable callus that produced plants. In a further study in 1988, Songstad, Duncan & Widholm in 10 *Plant Cell Reports* (1988), 7:262-265 reports several media additions that enhance regenerability of callus of two inbred lines. Other published reports also indicated that "nontraditional" tissues are capable of producing somatic embryogenesis and plant regeneration. K.P. Rao, et al., *Maize Genetics Cooperation Newsletter*, 60:64-65 (1986), refers to somatic embryogenesis from glume callus cultures and B.V. Conger, et al., 15 *Plant Cell Reports*, 6:345-347 (1987) indicates somatic embryogenesis from the tissue cultures of maize leaf segments. Thus, it is clear from the literature that the state of the art is such that these methods of obtaining plants are, and were, "conventional" in the sense that they are routinely used and have a very high rate of success.

Tissue culture of maize is described in European Patent Application, publication 20 160,390, incorporated herein by reference. Maize tissue culture procedures are also described in Green and Rhodes, "Plant Regeneration in Tissue Culture of Maize," *Maize for Biological Research* (Plant Molecular Biology Association, Charlottesville, Virginia 1982, at 367-372) and in Duncan, et al., "The Production of Callus Capable of Plant Regeneration from Immature Embryos of Numerous *Zea Mays* Genotypes," 165 *Planta* 25 322-332 (1985). Thus, another aspect of this invention is to provide cells which upon growth and differentiation produce maize plants having the physiological and morphological characteristics of inbred line PH48V.

 The utility of inbred maize line PH0KT also extends to crosses with other species. Commonly, suitable species will be of the family Gramineae, and especially of the genera *Zea*, *Tripsacum*, *Coix*, *Schlerachne*, *Polytoca*, *Chionachne*, and *Trilobachne*, of the tribe Maydeae. Potentially suitable for crosses with PH48V may be the various varieties of grain sorghum, *Sorghum bicolor* (L.) Moench.

Transformation of Maize

With the advent of molecular biological techniques that have allowed the

isolation and characterization of genes that encode specific protein products, scientists in the field of plant biology developed a strong interest in engineering the genome of plants to contain and express foreign genes, or additional, or modified versions of native or endogenous genes (perhaps driven by different promoters) in order to alter the traits of a plant in a specific manner. Such foreign, additional and/or modified genes are referred to herein collectively as "transgenes". Over the last fifteen to twenty years several methods for producing transgenic plants have been developed, and the present invention, in particular embodiments, also relates to transformed versions of the claimed inbred maize line PH48V.

Plant transformation involves the construction of an expression vector which will function in plant cells. Such a vector comprises DNA comprising a gene under control of or operatively linked to a regulatory element (for example, a promoter). The expression vector may contain one or more such operably linked gene/regulatory element combinations. The vector(s) may be in the form of a plasmid, and can be used, alone or in combination with other plasmids, to provide transformed maize plants, using transformation methods as described below to incorporate transgenes into the genetic material of the maize plant(s).

Expression Vectors For Maize Transformation

Marker Genes

Expression vectors include at least one genetic marker, operably linked to a regulatory element (a promoter, for example) that allows transformed cells containing the marker to be either recovered by negative selection, *i.e.* inhibiting growth of cells that do not contain the selectable marker gene, or by positive selection, *i.e.*, screening for the product encoded by the genetic marker. Many commonly used selectable marker genes for plant transformation are well known in the transformation arts, and include, for example, genes that code for enzymes that metabolically detoxify a selective chemical agent which may be an antibiotic or a herbicide, or genes that encode an altered target which is insensitive to the inhibitor. A few positive selection methods are also known in the art.

One commonly used selectable marker gene for plant transformation is the neomycin phosphotransferase II (*nptII*) gene, isolated from transposon Tn5, which when placed under the control of plant regulatory signals confers resistance to kanamycin. Fraley *et al.*, *Proc. Natl. Acad. Sci. U.S.A.*, 80: 4803 (1983). Another commonly used selectable marker gene is the hygromycin phosphotransferase gene which confers

resistance to the antibiotic hygromycin. Vanden Elzen *et al.*, *Plant Mol. Biol.*, 5: 299 (1985).

Additional selectable marker genes of bacterial origin that confer resistance to antibiotics include gentamycin acetyl transferase, streptomycin phosphotransferase, aminoglycoside- 3' -adenyl transferase, the bleomycin resistance determinant. Hayford *et al.*, *Plant Physiol.* 86: 1216 (1988), Jones *et al.*, *Mol. Gen. Genet.*, 210: 86 (1987), Svab *et al.*, *Plant Mol. Biol.*, 14: 197 (1990), Hille *et al.*, *Plant Mol. Biol.* 7: 171 (1986). Other selectable marker genes confer resistance to herbicides such as glyphosate, glufosinate or broxynil. Comai *et al.*, *Nature* 317: 741-744 (1985), Gordon-Kamm *et al.*, *Plant Cell* 2: 603-618 (1990) and Stalker *et al.*, *Science* 242: 419-423 (1988).

Other selectable marker genes for plant transformation are not of bacterial origin. These genes include, for example, mouse dihydrofolate reductase, plant 5 - enolpyruvylshikimate-3 -phosphate synthase and plant acetolactate synthase. Eichholtz *et al.*, *Somatic Cell Mol. Genet.* 13: 67 (1987), Shah *et al.*, *Science* 233: 478 (1986), Charest *et al.*, *Plant Cell Rep.* 8: 643 (1990).

Another class of marker genes for plant transformation require screening of presumptively transformed plant cells rather than direct genetic selection of transformed cells for resistance to a toxic substance such as an antibiotic. These genes are particularly useful to quantify or visualize the spatial pattern of expression of a gene in specific tissues and are frequently referred to as reporter genes because they can be fused to a gene or gene regulatory sequence for the investigation of gene expression. Commonly used genes for screening presumptively transformed cells include β -glucuronidase (*GUS*), β -galactosidase, luciferase and chloramphenicol acetyltransferase. Jefferson, R.A., *Plant Mol. Biol. Rep.* 5: 387 (1987), Teeri *et al.*, *EMBO J.* 8: 343 (1989), Koncz *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 84:131 (1987), De Block *et al.*, *EMBO J.* 3: 1681 (1984). Another approach to the identification of relatively rare transformation events has been use of a gene that encodes a dominant constitutive regulator of the *Zea mays* anthocyanin pigmentation pathway. Ludwig *et al.*, *Science* 247: 449 (1990).

Recently, *in vivo* methods for visualizing GUS activity that do not require destruction of plant tissue have been made available. Molecular Probes Publication 2908, Image Green™, p. 1-4 (1993) and Naleway *et al.*, *J. Cell Biol.* 115: 151a (1991). However, these *in vivo* methods for visualizing GUS activity have not proven useful for recovery of transformed cells because of low sensitivity, high fluorescent backgrounds,

and limitations associated with the use of luciferase genes as selectable markers.

More recently, a gene encoding Green Fluorescent Protein (GFP) has been utilized as a marker for gene expression in prokaryotic and eukaryotic cells. Chalfie *et al.*, *Science* 263: 802 (1994). GFP and mutants of GFP may be used as screenable markers.

Promoters

Genes included in expression vectors must be driven by a nucleotide sequence comprising a regulatory element, for example, a promoter. Several types of promoters are now well known in the transformation arts, as are other regulatory elements that can be used alone or in combination with promoters.

As used herein "promoter" includes reference to a region of DNA upstream from the start of transcription and involved in recognition and binding of RNA polymerase and other proteins to initiate transcription. A "plant promoter" is a promoter capable of initiating transcription in plant cells. Examples of promoters under developmental control include promoters that preferentially initiate transcription in certain tissues, such as leaves, roots, seeds, fibers, xylem vessels, tracheids, or sclerenchyma. Such promoters are referred to as "tissue-preferred". Promoters which initiate transcription only in certain tissues are referred to as "tissue-specific". A "cell type" specific promoter primarily drives expression in certain cell types in one or more organs, for example, vascular cells in roots or leaves. An "inducible" promoter is a promoter which is under environmental control. Examples of environmental conditions that may effect transcription by inducible promoters include anaerobic conditions or the presence of light. Tissue-specific, tissue-preferred, cell type specific, and inducible promoters constitute the class of "non-constitutive" promoters. A "constitutive" promoter is a promoter which is active under most environmental conditions.

A. Inducible Promoters

An inducible promoter is operably linked to a gene for expression in maize. Optionally, the inducible promoter is operably linked to a nucleotide sequence encoding a signal sequence which is operably linked to a gene for expression in maize. With an inducible promoter the rate of transcription increases in response to an inducing agent.

Any inducible promoter can be used in the instant invention. See Ward *et al.* *Plant Mol. Biol.* 22: 361-366 (1993). Exemplary inducible promoters include, but are not limited to, that from the ACEI system which responds to copper (Mett *et al.* *PNAS* 90: 4567-4571 (1993)); In2 gene from maize which responds to benzenesulfonamide

Science 23: 476-482 (1983) and Sengupta-Gopalan *et al.*, *Proc. Natl. Acad. Sci. USA* 82: 3320-3324 (1985)); a leaf-specific and light-induced promoter such as that from *cab* or *rubisco* (Simpson *et al.*, *EMBO J.* 4(11): 2723-2729 (1985) and Timko *et al.*, *Nature* 318: 579-582 (1985)); an anther-specific promoter such as that from *LAT52* (Twell *et al.*, *Mol. Gen. Genet.* 217: 240-245 (1989)); a pollen-specific promoter such as that from *Zm13* (Guerrero *et al.*, *Mol. Gen. Genet.* 224: 161-168 (1993)) or a microspore-preferred promoter such as that from *apg* (Twell *et al.*, *Sex. Plant Reprod.* 6: 217-224 (1993).

Signal Sequences For Targeting Proteins to Subcellular Compartments

Transport of protein produced by transgenes to a subcellular compartment such as the chloroplast, vacuole, peroxisome, glyoxysome, cell wall or mitochondrion, or for secretion into the apoplast, is accomplished by means of operably linking the nucleotide sequence encoding a signal sequence to the 5' and/or 3' region of a gene encoding the protein of interest. Targeting sequences at the 5' and/or 3' end of the structural gene may determine, during protein synthesis and processing, where the encoded protein is ultimately compartmentalized. The presence of a signal sequence directs a polypeptide to either an intracellular organelle or subcellular compartment or for secretion to the apoplast. Many signal sequences are known in the art. See, for example, Becker *et al.*, *Plant Mol. Biol.* 20: 49 (1992), Close, P.S., Master's Thesis, Iowa State University (1993), Knox, C., *et al.*, "Structure and Organization of Two Divergent Alpha-Amylase Genes From Barley", *Plant Mol. Biol.* 9: 3-17 (1987), Lerner *et al.*, *Plant Physiol.* 91: 124-129 (1989), Fontes *et al.*, *Plant Cell* 3: 483-496 (1991), Matsuoka *et al.*, *Proc. Natl. Acad. Sci.* 88: 834 (1991), Gould *et al.*, *J. Cell Biol* 108: 1657 (1989), Creissen *et al.*, *Plant J.* 2: 129 (1991), Kalderon, D., Robers, B., Richardson, W., and Smith A., "A short amino acid sequence able to specify nuclear location", *Cell* 39: 499-509 (1984), Stiefel, V., Ruiz-Avila, L., Raz R., Valles M., Gomez J., Pages M., Martinez-Izquierdo J., Ludevid M., Landale J., Nelson T., and Puigdomenech P., "Expression of a maize cell wall hydroxyproline-rich glycoprotein gene in early leaf and root vascular differentiation", *Plant Cell* 2: 785-793 (1990).

Foreign Protein Genes and Agronomic Genes

With transgenic plants according to the present invention, a foreign protein can be produced in commercial quantities. Thus, techniques for the selection and propagation of transformed plants, which are well understood in the art, yield a plurality of transgenic plants which are harvested in a conventional manner, and a foreign protein then can be extracted from a tissue of interest or from total biomass. Protein

extraction from plant biomass can be accomplished by known methods which are discussed, for example, by Heney and Orr, *Anal. Biochem.* 114: 92-6 (1981).

According to a preferred embodiment, the transgenic plant provided for commercial production of foreign protein is maize. In another preferred embodiment, the biomass of interest is seed. For the relatively small number of transgenic plants that show higher levels of expression, a genetic map can be generated, primarily via conventional Restriction Fragment Length Polymorphisms (RFLP), Polymerase Chain Reaction (PCR) analysis, and Simple Sequence Repeats (SSR) which identifies the approximate chromosomal location of the integrated DNA molecule. For exemplary methodologies in this regard, see Glick and Thompson, *METHODS IN PLANT MOLECULAR BIOLOGY AND BIOTECHNOLOGY* 269-284 (CRC Press, Boca Raton, 1993). Map information concerning chromosomal location is useful for proprietary protection of a subject transgenic plant. If unauthorized propagation is undertaken and crosses made with other germplasm, the map of the integration region can be compared to similar maps for suspect plants, to determine if the latter have a common parentage with the subject plant. Map comparisons would involve hybridizations, RFLP, PCR, SSR and sequencing, all of which are conventional techniques.

Likewise, by means of the present invention, agronomic genes can be expressed in transformed plants. More particularly, plants can be genetically engineered to express various phenotypes of agronomic interest. Exemplary genes implicated in this regard include, but are not limited to, those categorized below.

1. Genes That Confer Resistance To Pests or Disease And That Encode:

(A) Plant disease resistance genes. Plant defenses are often activated by specific interaction between the product of a disease resistance gene (R) in the plant and the product of a corresponding avirulence (Avr) gene in the pathogen. A plant variety can be transformed with cloned resistance gene to engineer plants that are resistant to specific pathogen strains. See, for example Jones et al., *Science* 266: 789 (1994) (cloning of the tomato Cf-9 gene for resistance to *Cladosporium fulvum*); Martin et al., *Science* 262: 1432 (1993) (tomato *Pto* gene for resistance to *Pseudomonas syringae* pv. tomato encodes a protein kinase); Mindrinos et al., *Cell* 78: 1089 (1994) (*Arabidopsis RSP2* gene for resistance to *Pseudomonas syringae*).

(B) A *Bacillus thuringiensis* protein, a derivative thereof or a synthetic polypeptide modeled thereon. See, for example, Geiser et al., *Gene* 48: 109 (1986), who disclose the cloning and nucleotide sequence of a *Bt* δ -endotoxin gene. Moreover,

licheniformis α -amylase), Elliot *et al.*, *Plant Molec. Biol.* 21: 515 (1993) (nucleotide sequences of tomato invertase genes), Sogaard *et al.*, *J. Biol. Chem.* 268: 22480 (1993) (site-directed mutagenesis of barley α -amylase gene), and Fisher *et al.*, *Plant Physiol.* 102: 1045 (1993) (maize endosperm starch branching enzyme II).

5

Methods for Maize Transformation

Numerous methods for plant transformation have been developed, including biological and physical, plant transformation protocols. See, for example, Miki *et al.*, "Procedures for Introducing Foreign DNA into Plants" in *Methods in Plant Molecular Biology and Biotechnology*, Glick, B.R. and Thompson, J.E. Eds. (CRC Press, Inc., Boca Raton, 1993) pages 67-88. In addition, expression vectors and *in vitro* culture methods for plant cell or tissue transformation and regeneration of plants are available. See, for example, Gruber *et al.*, "Vectors for Plant Transformation" in *Methods in Plant Molecular Biology and Biotechnology*, Glick, B.R. and Thompson, J.E. Eds. (CRC Press, Inc., Boca Raton, 1993) pages 89-119.

A. *Agrobacterium*-mediated Transformation

One method for introducing an expression vector into plants is based on the natural transformation system of *Agrobacterium*. See, for example, Horsch *et al.*, *Science* 227: 1229 (1985). *A. tumefaciens* and *A. rhizogenes* are plant pathogenic soil bacteria which genetically transform plant cells. The Ti and Ri plasmids of *A. tumefaciens* and *A. rhizogenes*, respectively, carry genes responsible for genetic transformation of the plant. See, for example, Kado, C.I., *Crit. Rev. Plant. Sci.* 10: 1 (1991). Descriptions of *Agrobacterium* vector systems and methods for *Agrobacterium*-mediated gene transfer are provided by Gruber *et al.*, *supra*, Miki *et al.*, *supra*, and Moloney *et al.*, *Plant Cell Reports* 8: 238 (1989). See also, U.S. Patent No. 5,591,616, issued Jan. 7, 1997.

B. Direct Gene Transfer

Despite the fact the host range for *Agrobacterium*-mediated transformation is broad, some major cereal crop species and gymnosperms have generally been recalcitrant to this mode of gene transfer, even though some success has recently been achieved in rice and maize. Hiei *et al.*, *The Plant Journal* 6: 271-282 (1994); U.S. Patent No. 5,591,616, issued Jan. 7, 1997. Several methods of plant transformation, collectively referred to as direct gene transfer, have been developed as an alternative to *Agrobacterium*-mediated transformation.

A generally applicable method of plant transformation is microprojectile-mediated transformation wherein DNA is carried on the surface of microprojectiles measuring 1 to 4 μm . The expression vector is introduced into plant tissues with a biolistic device that accelerates the microprojectiles to speeds of 300 to 600 m/s which is sufficient to penetrate plant cell walls and membranes. Sanford *et al.*, *Part. Sci. Technol.* 5: 27 (1987), Sanford, J.C., *Trends Biotech.* 6: 299 (1988), Klein *et al.*, *Bio/Technology* 6: 559-563 (1988), Sanford, J.C., *Physiol Plant* 79: 206 (1990), Klein *et al.*, *Biotechnology* 10: 268 (1992). In maize, several target tissues can be bombarded with DNA-coated microprojectiles in order to produce transgenic plants, including, for example, callus (Type I or Type II), immature embryos, and meristematic tissue.

Another method for physical delivery of DNA to plants is sonication of target cells. Zhang *et al.*, *Bio/Technology* 9: 996 (1991). Alternatively, liposome or spheroplast fusion have been used to introduce expression vectors into plants. Deshayes *et al.*, *EMBO J.*, 4: 2731 (1985), Christou *et al.*, *Proc Natl. Acad. Sci. U.S.A.* 84: 3962 (1987). Direct uptake of DNA into protoplasts using CaCl_2 precipitation, polyvinyl alcohol or poly-L-ornithine have also been reported. Hain *et al.*, *Mol. Gen. Genet.* 199: 161 (1985) and Draper *et al.*, *Plant Cell Physiol.* 23: 451 (1982). Electroporation of protoplasts and whole cells and tissues have also been described. Donn *et al.*, In Abstracts of VIIth International Congress on Plant Cell and Tissue Culture IAPTC, A2-38, p 53 (1990); D'Halluin *et al.*, *Plant Cell* 4: 1495-1505 (1992) and Spencer *et al.*, *Plant Mol. Biol.* 24: 51-61 (1994).

Following transformation of maize target tissues, expression of the above-described selectable marker genes allows for preferential selection of transformed cells, tissues and/or plants, using regeneration and selection methods now well known in the art. For example, transformed maize immature embryos.

The foregoing methods for transformation would typically be used for producing transgenic inbred lines. Transgenic inbred lines could then be crossed, with another (non-transformed or transformed) inbred line, in order to produce a transgenic hybrid maize plant. Alternatively, a genetic trait which has been engineered into a particular maize line using the foregoing transformation techniques could be moved into another line using traditional backcrossing techniques that are well known in the plant breeding arts. For example, a backcrossing approach could be used to move an engineered trait from a public, non-elite line into an elite line, or from a hybrid maize plant containing a foreign gene in its genome into a line or lines which do not contain that gene. As used

Inbred Comparisons

The results in Table 2A compare inbred PH48V to inbred PH00M. The results show inbred PH48V demonstrates above average yields. Inbred PH48V presents a significantly taller plant than inbred PH00M. Inbred PH48V shows very good resistance to root lodging and to stalk lodging and has a high staygreen score. Inbred PH48V shows excellent resistance to Gray Leaf Spot.

The results in Table 2B compare inbred PH48V to inbred PHKV1. The results show inbred PH48V demonstrates above average yields. Inbred PH48V presents a significantly taller plant than inbred PHKV1. Inbred PH48V shows very good resistance to root lodging and to stalk lodging and has a significantly higher staygreen score than inbred PHKV1. Inbred PH48V shows strong resistance to Gray Leaf Spot and shows significantly better resistance than inbred PHKV1. Inbred PH48V shows strong resistance to Northern Leaf Blight and Southern Leaf Blight.

The results in Table 2C compare inbred PH48V to inbred PH07D. The results show inbred PH48V demonstrates above average yields. Inbred PH48V presents a significantly taller plant with significantly higher ear placement than inbred PH07D. Inbred PH48V shows very good resistance to root lodging and to stalk lodging and has a significantly higher staygreen score than inbred PH07D. Inbred PH48V shows strong resistance to Gray Leaf Spot and shows significantly better resistance than inbred PH07D. Inbred PH48V shows strong resistance to Northern Leaf Blight and Southern Leaf Blight.

Inbred By Tester Comparisons

The results in Table 3 compare the inbred PH48V and inbred PH07D, when each inbred is crossed to the same tester lines. The PH48V hybrid demonstrates above average and significantly higher yields than the PH07D hybrid. The PH48V hybrid presents a plant with higher than average ear placement. The PH48V hybrid shows above average resistance to root lodging and stalk lodging.

Hybrid Comparisons

The results in Table 4A compare inbred PH48V crossed to inbred PH0KT and inbred PHKV1 crossed to PHN46. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid presents higher than average ear placement and a

significantly taller plant than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid shows significantly better staygreen scores than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to stalk lodging and brittle stalk than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PHKV1/PHN46 hybrid.

The results in Table 4B compare inbred PH48V crossed to inbred PH0KT and inbred PHW52 crossed to PHK46. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields with significantly higher test weight of grain than the PHW52/PHK46 hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PHW52/PHK46 hybrid. The PH48V/PH0KT hybrid shows above average staygreen scores and demonstrates above average resistance to root lodging. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to stalk lodging and brittle stalk than the PHW52/PHK46 hybrid. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PHW52/PHK46 hybrid.

The results in Table 4C compare inbred PH48V crossed to inbred PH0KT and inbred PH6M1 crossed to PH24M. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PH6M1/PH24M hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PH6M1/PH24M hybrid. The PH48V/PH0KT hybrid shows above average staygreen scores. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to stalk lodging than the PH6M1/PH24M hybrid. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PH6M1/PH24M hybrid and above average resistance to Northern Leaf Blight.

The results in Table 4D compare inbred PH48V crossed to inbred PH0KT and inbred PH07D crossed to PH0V0. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PH07D/PH0V0 hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PH07D/PH0V0 hybrid. The PH48V/PH0KT hybrid shows above average staygreen scores. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to brittle stalk than the PH07D/PH0V0 hybrid,

and shows above average resistance to stalk lodging. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Southern Leaf Blight than the PH07D/PH0V0 hybrid and above average resistance to Gray Leaf Spot and Northern Leaf Blight.

- 5 The results in Table 4E compare inbred PH48V crossed to inbred PH0KT and inbred PH07D crossed to PH24M. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PH07D/PH24M hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PH07D/PH24M hybrid. The PH48V/PH0KT hybrid shows
- 10 above average staygreen scores and demonstrates above average resistance to stalk lodging. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PH07D/PH24M hybrid and above average resistance to Northern Leaf Blight.

TABLE 2A
PAIRED INBRED COMPARISON REPORT

VARIETY #1 = PH48V
VARIETY #2 = PHOOM

5				BU	BU		TST	EGR	EST	TIL	GDU	GDU
				ACR	ACR	MST	WT	WTH	CNT	LER	SHD	SLK
				ABS	%MN	ABS	ABS	ABS	ABS	ABS	ABS	ABS
10	TOTAL SUM	1		71.2	105	26.1	56.0	4.8	38.5	2.4	148.7	154.6
		2		57.6	91	26.1	54.5	4.4	35.5	1.3	147.7	154.8
		LOCS		12	12	14	2	9	22	13	20	20
		REPS		13	13	15	2	10	23	13	20	20
15		DIFF		13.5	14	0.0	1.5	0.4	3.0	1.1	1.0	0.3
		PR > T		.111	.403	.999	.066*	.211	.018+	.276	.347	.838
20				POL	TAS	PLT	EAR	RT	STA	STK	BRT	SCT
				SC	SZ	HT	HT	LDG	GRN	LDG	STK	GRN
				ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
25	TOTAL SUM	1		5.7	5.3	92.7	39.5	98.7	7.3	100.0	100.0	7.0
		2		4.7	4.6	85.7	34.0	100.0	6.5	93.1	100.0	3.0
		LOCS		3	13	6	4	3	4	2	1	1
		REPS		3	13	6	4	3	4	2	1	1
		DIFF		1.0	0.7	7.0	5.5	1.3	0.8	6.9	0.0	4.0
30		PR > T		.000#	.056*	.001#	.086*	.423	.486	.500		
35				EAR	TEX	EAR	BAR	GLF	STW	COM	CLD	CLD
				SZ	EAR	MLD	PLT	SPT	WLT	RST	TST	TST
				ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	%MN
40	TOTAL SUM	1		5.0	6.0	7.0	98.4	9.0	8.0	5.3	78.0	108
		2		4.0	5.5	4.5	92.2	8.5	8.0	5.5	63.7	87
		LOCS		1	2	2	11	2	1	4	3	3
		REPS		1	2	2	11	2	1	4	3	3
		DIFF		1.0	0.5	2.5	6.2	0.5	0.0	0.3	14.3	21
45		PR > T			.500	.344	.060*	.500		.391	.246	.258
50				KSZ								
				DCD								
				ABS								
55	TOTAL SUM	1		5.7								
		2		5.3								
		LOCS		3								
		REPS		3								
		DIFF		0.3								
60		PR > T		.945								
	* = 10% SIG											
	+ = 5% SIG											
	# = 1% SIG											

TABLE 2B
 PAIRED INBRED COMPARISON REPORT

VARIETY #1 = PH48V
 VARIETY #2 = PHKV1

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		BU	BU		TST	EGR	EST	TIL	GDU	GDU
		ACR	ACR	MST	WT	WTH	CNT	LER	SHD	SLK
		ABS	%MN	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL SUM	1	72.5	109	26.6	54.8	5.0	30.9	2.7	151.5	157.6
	2	64.5	96	23.7	55.1	4.8	31.2	1.5	155.8	157.5
	LOCS	35	35	39	9	26	47	40	56	54
	REPS	37	37	41	10	27	48	41	56	54
	DIFF	8.0	13	2.9	0.3	0.1	0.3	1.2	4.2	0.1
	PR > T	.066*	.061*	.000#	.472	.622	.594	.068*	.000#	.883

		POL	TAS	PLT	EAR	RT	STA	STK	BRT	SCT
		SC	SZ	HT	HT	LDG	GRN	LDG	STK	GRN
		ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL SUM	1	5.7	5.4	95.7	41.2	96.3	7.8	100.0	100.0	6.8
	2	4.3	2.7	89.2	38.7	89.7	4.7	80.8	100.0	6.4
	LOCS	3	36	25	13	5	11	2	3	5
	REPS	3	36	25	13	5	12	2	3	5
	DIFF	1.3	2.8	6.5	2.5	6.5	3.1	19.2	0.0	0.4
	PR > T	.057*	.000#	.000#	.018+	.257	.001#	.500	.999	.374

		EAR	TEX	EAR	BAR	GLF	NLF	SLF	STW	ANT
		SZ	EAR	MLD	PLT	SPT	BLT	BLT	WLT	ROT
		ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL SUM	1	5.0	6.0	7.6	94.5	7.9	6.5	6.0	7.3	4.5
	2	6.0	6.0	7.6	89.7	2.6	3.5	7.0	5.5	4.5
	LOCS	1	2	5	36	9	1	1	3	2
	REPS	1	2	5	36	12	2	2	4	3
	DIFF	1.0	0.0	0.0	4.9	5.4	3.0	1.0	1.8	0.0
	PR > T		.999	.999	.062*	.000#			.008#	.999

		HD	MDM	FUS	DIP	COM	CLD	CLD	KSZ
		SMT	CPX	ERS	ERS	RST	TST	TST	DCD
		ABS	ABS	ABS	ABS	ABS	ABS	%MN	ABS
TOTAL SUM	1	100.0	4.0	6.7	1.5	5.3	85.2	107	3.6
	2	100.0	4.0	7.2	2.5	1.3	83.3	105	1.8
	LOCS	1	1	6	1	4	15	15	15
	REPS	2	2	7	2	4	15	15	15
	DIFF	0.0	0.0	0.5	1.0	4.0	1.9	2	1.8
	PR > T			.296		.011+	.670	.751	.039+

* = 10% SIG
 + = 5% SIG
 # = 1% SIG

[illegible]

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* = 10% SIG

+ = 5% SIG

= 1% SIG

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TABLE 3

Average Inbred By Tester Performance Comparing PH48V To PH07D Crossed
To The Same Inbred Testers And Grown In The Same Experiments.

		BU ACR ABS	BU ACR %MN	MST %MN	EGR WTH %MN	EST CNT %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN
TOTAL SUM	REPS	21	21	21	5	8	20	9	9	4	10
	LOCS	21	21	21	5	8	20	9	9	4	10
	PH48V	170	114	102	85	105	104	101	110	103	107
	PH07D	156	105	93	93	97	100	99	100	103	127
	DIFF	14	9	9	7	8	3	2	9	0	20
	PR > T	0.00	0.00	0.00	0.50	0.01	0.08	0.34	0.01	0.99	0.09
		STK LDG %MN	SLF BLT ABS	COM RST ABS	SOU RST ABS	DRP EAR %MN	GLF SPT ABS				
TOTAL SUM	REPS	13	1	3	1	1	1				
	LOCS	13	1	3	1	1	1				
	PH48V	104	7	7	4	100	7				
	PH07D	105	6	6	6	100	5				
	DIFF	1	1	1	2	0	2				
	PR > T	0.56		0.23		0.99					

*PR > T values are valid only for comparisons with Locs >= 10.

TABLE 4A
INBREDS IN HYBRID COMBINATION REPORT

		VARIETY #1 = PH48V/PH0KT								
		VARIETY #2 = PHKV1/PHN46								
		PRM	SHD	BU	BU	MST	TST	EGR	EST	GDU
		ABS	ABS	ABS	%MN	%MN	WT	WTH	CNT	SHD
		ABS	ABS	ABS	%MN	%MN	ABS	%MN	%MN	%MN
TOTAL SUM	1	119	116	173.7	106	105	57.0	92	103	100
	2	115	117	166.9	101	94	57.2	92	104	103
	LOCS	7	6	143	143	143	58	27	30	26
	REPS	7	6	165	165	165	59	30	34	29
	DIFF	4	1	6.8	4	11	0.3	0	1	3
	PR > T	.002#	.007#	.001#	.001#	.000#	.252	.999	.733	.000#
		GDU	STK	PLT	EAR	RT	STA	STK	BRT	DRP
		SLK	CNT	HT	HT	LDG	GRN	LDG	STK	EAR
		%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN
TOTAL SUM	1	101	101	104	108	99	109	102	108	100
	2	102	102	100	107	103	86	97	98	100
	LOCS	17	190	51	50	31	47	55	14	9
	REPS	20	245	62	60	34	52	59	19	11
	DIFF	2	1	4	2	4	23	6	11	0
	PR > T	.014+	.183	.000#	.195	.027+	.000#	.002#	.037+	.999
		GLF	NLF	SLF	STW	ANT	HD	MDM	FUS	
		SPT	BLT	BLT	WLT	ROT	SMT	CLN	CPX	ERS
		ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL SUM	1	6.6	5.8	7.4	7.0	5.0	96.9	6.3	3.0	4.9
	2	2.7	5.8	3.6	5.8	2.9	79.6	6.8	3.0	5.3
	LOCS	8	5	7	5	6	2	2	1	9
	REPS	12	7	10	5	10	4	4	2	12
	DIFF	3.9	0.0	3.8	1.2	2.1	17.3	0.5	0.0	0.4
	PR > T	.000#	.999	.000#	.235	.021+	.500	.500		.548
		* = 10% SIG								
		+ = 5% SIG								
		# = 1% SIG								

TABLE 4A (cont.)
INBREDS IN HYBRID COMBINATION REPORT

		VARIETY #1 = PH48V/PHOKT VARIETY #2 = PHKV1/PHN46									
		DIP	COM	SOU	ECB	ECB	HSK	HSK	OIL	PRO	
		ERS	RST	RST	1LF	2SC	CVR	CVR	T	T	
		ABS	ABS	ABS	ABS	ABS	ABS	%MN	ABS	ABS	
TOTAL	SUM	1	2.8	6.3	4.0	7.5	4.5	5.0	90	4.4	9.0
		2	3.3	3.4	4.0	6.3	3.5	5.0	90	4.0	9.2
	LOCS	2	10	1	3	1	16	16	8	8	8
	REPS	4	11	1	6	2	16	16	8	8	8
	DIFF	0.5	2.9	0.0	1.2	1.0	0.0	0	0.3	0.2	0.2
	PR > T	.500	.000#		.020+		.999	.999	.149	.639	

STR											
T											
ABS											

TOTAL	SUM	1	72.3								
		2	72.6								
	LOCS	8									
	REPS	8									
	DIFF	0.3									
	PR > T	.221									

* = 10% SIG											
+ = 5% SIG											
# = 1% SIG											

0040066 012100

TABLE 4B
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PHOKT
VARIETY #2 = PHW52/PHK46

5	VARIETY #2 = PHW52/PHK46										
			PRM	PRM	BU	BU	TST	EGR	EST	GDU	
			ABS	SHD	ACR	ACR	WT	WTH	CNT	SHD	
				ABS	ABS	%MN	%MN	%MN	%MN	%MN	
10	TOTAL SUM	1	120	115	176.6	106	106	57.0	91	103	100
		2	118	116	167.1	100	102	56.0	99	100	101
	LOCS		5	5	125	125	125	66	25	33	23
	REPS		5	5	140	140	140	67	27	37	25
15	DIFF		1	0	9.5	6	4	1.0	7	3	1
	PR > T		.006#	.999	.000#	.000#	.000#	.000#	.091*	.098*	.009#
20											
			GDU	STK	PLT	EAR	RT	STA	STK	BRT	DRP
			SLK	CNT	HT	HT	LDG	GRN	LDG	STK	EAR
			%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN
25	TOTAL SUM	1	100	101	104	108	101	111	103	108	100
		2	101	98	100	101	98	107	97	90	100
	LOCS		15	160	40	40	13	40	38	13	9
	REPS		17	197	42	42	13	43	40	17	11
	DIFF		1	2	4	7	3	4	5	18	0
30	PR > T		.203	.007#	.000#	.000#	.384	.459	.010+	.023+	.999
35											
			GLF	NLF	SLF	STW	ANT	HD	MDM	FUS	
			SPT	BLT	BLT	WLT	ROT	SMT	CLN	CPX	ERS
			ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
40	TOTAL SUM	1	6.6	5.8	7.4	7.0	5.0	96.9	6.3	3.0	5.1
		2	4.4	5.3	6.5	4.2	4.1	72.9	6.3	3.0	6.2
	LOCS		8	4	7	5	6	2	2	1	9
	REPS		12	6	10	5	10	4	4	2	12
	DIFF		2.2	0.5	0.9	2.8	0.9	24.0	0.0	0.0	1.1
45	PR > T		.000#	.423	.007#	.009#	.186	.426	.999		.159
	* = 10% SIG										
	+ = 5% SIG										
	# = 1% SIG										

TABLE 4B (cont.)
INBREDS IN HYBRID COMBINATION REPORT

		VARIETY #1 = PH48V/PHOKT VARIETY #2 = PHW52/PHK46									
5			DIP ERS ABS	COM RST ABS	SOU RST ABS	ECB 1LF ABS	ECB 2SC ABS	HSK CVR ABS	HSK CVR %MN	OIL T ABS	PRO T ABS
10	TOTAL SUM	1	2.8	6.9	4.0	7.5	4.5	5.1	92	4.3	9.2
		2	3.0	6.5	5.0	6.0	4.0	5.1	90	4.3	8.8
	LOCS		2	5	1	3	1	17	17	6	6
	REPS		4	6	1	6	2	17	17	6	6
15	DIFF		0.3	0.4	1.0	1.5	0.5	0.0	2	0.1	0.5
	PR > T		.500	.374		.122		.999	.776	.648	.422

20			STR T ABS								
25	TOTAL SUM	1	72.2								
		2	72.5								
	LOCS		6	.							
	REPS		6	.							
30	DIFF		0.3								
	PR > T		.492								

	* = 10% SIG										
	+ = 5% SIG										
	# = 1% SIG										

TABLE 4C
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PH0KT
VARIETY #2 = PH6M1/PH24M

5	VARIETY #2 = PH6M1/PH24M											
			PRM	PRM	BU	BU		TST	EGR	EST	GDU	
			ABS	SHD	ACR	ACR	MST	WT	WTH	CNT	SHD	
10				ABS	ABS	%MN	%MN	ABS	%MN	%MN	%MN	
	TOTAL	SUM	1	119	115	176.8	104	106	57.4	95	104	100
			2	115	116	163.6	96	94	58.9	105	101	102
			LOCS	4	3	78	78	35	14	17	17	17
			REPS	4	3	93	93	36	16	21	19	19
15			DIFF	4	1	13.2	8	11	1.5	10	3	2
			PR > T	.002#	.022+	.000#	.000#	.000#	.051*	.134	.001#	
20												
			GDU	STK	PLT	EAR	RT	STA	STK	BRT	DRP	
			SLK	CNT	HT	HT	LDG	GRN	LDG	STK	EAR	
			%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	
25	TOTAL	SUM	1	100	101	105	109	100	112	102	108	100
			2	103	100	102	100	102	96	98	100	100
			LOCS	12	114	29	28	7	25	22	13	8
			REPS	14	157	36	34	7	28	24	17	10
			DIFF	2	1	3	9	2	16	4	8	0
30			PR > T	.003#	.337	.011+	.000#	.326	.064*	.004#	.278	.999
35												
			GLF	NLF	SLF	STW	ANT	HD		MDM	FUS	
			SPT	BLT	BLT	WLT	ROT	SMT	CLN	CPX	ERS	
			ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	
40	TOTAL	SUM	1	6.5	5.8	7.6	7.0	5.0	96.9	6.3	3.0	5.3
			2	3.9	4.4	6.7	6.0	4.4	77.1	6.5	3.0	5.3
			LOCS	7	4	5	5	6	2	2	1	8
			REPS	11	6	8	5	10	4	4	2	10
			DIFF	2.6	1.4	0.9	1.0	0.6	19.8	0.3	0.0	0.0
45			PR > T	.003#	.062*	.001#	.142	.363	.500	.500		.999
50												
			DIP	COM	HSK	HSK	OIL	PRO	STR			
			ERS	RST	CVR	CVR	T	T	T			
			ABS	ABS	ABS	%MN	ABS	ABS	ABS			
55	TOTAL	SUM	1	2.8	7.3	5.1	93	4.4	9.0	72.3		
			2	3.5	6.3	5.5	101	4.2	9.7	71.6		
			LOCS	2	2	8	8	8	8	8		
			REPS	4	3	8	8	8	8	8		
			DIFF	0.8	1.0	0.4	8	0.2	0.6	0.8		
60			PR > T	.205	.000#	.351	.351	.215	.232	.004#		
	* = 10% SIG											
	+ = 5% SIG											
	# = 1% SIG											

TABLE 4D
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PH0KT
VARIETY #2 = PH07D/PH0V0

5	VARIETY #2 = PH07D/PH0V0											
			PRM	PRM	BU	BU		TST	EGR	EST	GDU	
			ABS	SHD	ACR	ACR	MST	WT	WTH	CNT	SHD	
10				ABS	ABS	%MN	%MN	ABS	%MN	%MN	%MN	
	TOTAL	SUM	1	119	116	169.3	104	105	57.2	96	104	100
			2	118	116	155.1	95	103	57.5	104	102	101
		LOCS		5	4	94	94	34	18	17	20	
		REPS		5	4	108	108	35	21	21	23	
15		DIFF		1	0	14.2	9	3	0.3	8	3	1
		PR > T		.327	.999	.000#	.000#	.002#	.224	.026+	.202	.203
20												
			GDU	STK	PLT	EAR	RT	STA	STK	BRT	DRP	
			SLK	CNT	HT	HT	LDG	GRN	LDG	STK	EAR	
			%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	
25	TOTAL	SUM	1	101	101	105	108	98	110	102	108	100
			2	101	103	101	101	100	108	99	90	100
		LOCS		14	136	37	36	25	32	38	14	8
		REPS		17	182	48	46	28	37	42	19	10
		DIFF		0	1	3	7	2	2	3	18	0
30		PR > T		.999	.085*	.001#	.000#	.296	.802	.177	.030+	.999
35												
			GLF	NLF	SLF	STW	ANT	HD		MDM	FUS	
			SPT	BLT	BLT	WLT	ROT	SMT	CLN	CPX	ERS	
			ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	
40	TOTAL	SUM	1	6.5	5.8	7.6	7.0	5.0	96.9	6.3	3.0	5.0
			2	5.9	6.4	5.9	6.4	4.7	100.0	6.5	3.0	6.4
		LOCS		7	5	5	5	6	2	2	1	9
		REPS		11	7	8	5	10	4	4	2	11
		DIFF		0.6	0.6	1.7	0.6	0.3	3.1	0.3	0.0	1.4
45		PR > T		.122	.109	.010+	.208	.655	.500	.500		.069*
50												
			DIP	COM	HSK	HSK	OIL	PRO	STR			
			ERS	RST	CVR	CVR	T	T	T			
			ABS	ABS	ABS	%MN	ABS	ABS	ABS			
55	TOTAL	SUM	1	3.3	6.1	5.1	93	4.3	9.4	72.1		
			2	3.5	5.2	6.1	111	4.2	9.1	71.8		
		LOCS		2	7	8	8	7	7	7		
		REPS		3	8	8	8	7	7	7		
		DIFF		0.3	0.9	1.0	19	0.1	0.4	0.3		
60		PR > T		.874	.045+	.033+	.034+	.687	.551	.293		
	* = 10% SIG											
	+ = 5% SIG											
	# = 1% SIG											

TABLE 4E
INBREDS IN HYBRID COMBINATION REPORT

		VARIETY #1 = PH48V/PHOKT VARIETY #2 = PH07D/PH24M									
			PRM ABS	PRM SHD ABS	BU ACR ABS	BU ACR %MN	MST %MN	TST WT ABS	EGR WTH %MN	EST CNT %MN	GDU SHD %MN
5	TOTAL SUM	1	119	115	169.9	105	105	57.4	93	103	100
10		2	114	115	160.8	100	94	59.3	103	102	101
	LOCS		6	4	119	119	120	49	21	29	20
	REPS		6	4	134	134	135	50	23	33	22
15	DIFF		5	0	9.2	6	11	1.9	10	1	1
	PR > T		.000#	.999	.000#	.000#	.000#	.000#	.042+	.449	.161
			GDU SLK %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN	STK LDG %MN	BRT STK %MN	DRP EAR %MN
20	TOTAL SUM	1	100	101	104	109	99	112	102	108	100
25		2	101	102	100	96	102	102	104	104	100
	LOCS		12	159	42	41	25	35	49	13	9
	REPS		14	207	52	50	25	38	51	17	11
30	DIFF		0	0	4	13	3	10	2	4	0
	PR > T		.999	.999	.000#	.000#	.089*	.097*	.042+	.406	.999
			GLF SPT ABS	NLF BLT ABS	SLF BLT ABS	STW WLT ABS	ANT ROT ABS	HD SMT ABS	CLN ABS	MDM CPX ABS	FUS ERS ABS
35	TOTAL SUM	1	6.6	5.8	7.4	7.0	5.0	96.9	6.3	3.0	4.8
40		2	5.5	5.5	5.3	6.0	4.9	93.8	6.5	3.0	4.8
	LOCS		8	4	6	5	6	2	2	1	8
	REPS		12	6	8	5	10	4	4	2	11
45	DIFF		1.1	0.3	2.1	1.0	0.1	3.1	0.3	0.0	0.1
	PR > T		.046+	.638	.000#	.230	.910	.500	.500		.945
			DIP ERS ABS	COM RST ABS	SOU RST ABS	HSK CVR ABS	HSK CVR %MN	OIL T ABS	PRO T ABS	STR T ABS	
50	TOTAL SUM	1	3.3	6.2	4.0	5.1	92	4.3	9.3	71.8	
55		2	3.0	6.0	4.0	5.2	94	4.0	9.7	70.9	
	LOCS		2	10	1	15	15	3	3	3	
	REPS		3	10	1	15	15	3	3	3	
60	DIFF		0.3	0.2	0.0	0.1	2	0.2	0.4	0.9	
	PR > T		.795	.443		.653	.673	.113	.414	.058*	
		* = 10% SIG + = 5% SIG # = 1% SIG									

Deposits

A deposit of the seed of inbred PH48V is and has been maintained by Pioneer Hi-Bred International, Inc., 800 Capital Square, 400 Locust Street, Des Moines, Iowa 50309-2340, since prior to the filing date of this application. Access to this deposit will
5 be available during the pendency of the application to the Commissioner of Patents and Trademarks and person determined by the Commissioner to be entitled thereto upon request. Upon allowance of any claims in the application, the Applicant(s) will make available to the public without restriction a deposit of at least 2500 seeds of inbred PH48V with the American Type Culture Collection (ATCC), Manassas, Virginia, 20110. The seeds deposited with the ATCC will be taken from the same deposit maintained at Pioneer Hi-Bred and described above. Additionally, Applicant(s) will meet all the requirements of 37 C.F.R. §1.801 - 1.809, including providing an indication of the viability of the sample when the deposit is made. This deposit of Inbred Maize Line PH48V will be maintained in the ATCC Depository, which is a public depository, for a
15 period of 30 years, or 5 years after the most recent request, or for the enforceable life of the patent, whichever is longer, and will be replaced if it ever becomes nonviable during that period. Applicant will impose no restrictions on the availability of the deposited material from the ATCC; however, Applicant has no authority to waive any restrictions imposed by law on the transfer of biological material or its transportation in commerce. Applicant does not waive any infringement of its rights granted under this patent or
20 under the Plant Variety Protection Act (7 USC 2321 et seq.).

The foregoing invention has been described in detail by way of illustration and example for purposes of clarity and understanding. However, it will be obvious that certain changes and modifications such as single gene conversions and mutations,
25 somoclonal variants, variant individuals selected from large populations of the plants of the instant inbred and the like may be practiced within the scope of the invention, as limited only by the scope of the appended claims.